

Appl. No. : 10/063,715
Filed : May 8, 2002

AMENDMENTS TO THE CLAIMS

1. (Currently Amended) An isolated nucleic acid having at least 80% nucleic acid sequence identity to:
 - (a) a nucleic acid sequence encoding the polypeptide ~~of shown in Figure 88~~ (SEQ ID NO:88);
 - (b) a nucleic acid sequence encoding the polypeptide ~~of shown in Figure 88~~ (SEQ ID NO:88), lacking its associated signal peptide;
 - (c) ~~a nucleic acid sequence encoding the extracellular domain of the polypeptide of shown in Figure 88 (SEQ ID NO:88);~~
 - (d) ~~a nucleic acid sequence encoding the extracellular domain of the polypeptide of shown in Figure 88 (SEQ ID NO:88) lacking its associated signal peptide;~~
 - (e) ~~(c) the nucleic acid sequence of shown in Figure 87 (SEQ ID NO:87);~~
 - (f) (d) the full-length coding sequence of the nucleic acid sequence ~~of shown in Figure 87~~ (SEQ ID NO:87); or
 - (g) (e) the full-length coding sequence of the cDNA deposited under ATCC accession number 203159;
wherein said isolated nucleic acid is more highly expressed in normal lung tissue compared to lung tumor, or wherein said isolated nucleic acid encodes a polypeptide that is more highly expressed in normal lung tissue compared to lung tumor.
2. (Currently Amended) The isolated nucleic acid of Claim 1 having at least 85% nucleic acid sequence identity to:
 - (a) a nucleic acid sequence encoding the polypeptide ~~of shown in Figure 88~~ (SEQ ID NO:88);
 - (b) a nucleic acid sequence encoding the polypeptide ~~of shown in Figure 88~~ (SEQ ID NO:88), lacking its associated signal peptide;
 - (c) ~~a nucleic acid sequence encoding the extracellular domain of the polypeptide of shown in Figure 88 (SEQ ID NO:88);~~
 - (d) ~~a nucleic acid sequence encoding the extracellular domain of the polypeptide of shown in Figure 88 (SEQ ID NO:88) lacking its associated signal peptide;~~
 - (e) ~~(c) the nucleic acid sequence of shown in Figure 87 (SEQ ID NO:87);~~

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(f) (d) the full-length coding sequence of the nucleic acid sequence of shown in Figure 87 (SEQ ID NO:87); or

(g) (e) the full-length coding sequence of the cDNA deposited under ATCC accession number 203159;

wherein said isolated nucleic acid is more highly expressed in normal lung tissue compared to lung tumor, or wherein said isolated nucleic acid encodes a polypeptide that is more highly expressed in normal lung tissue compared to lung tumor.

3. (Currently Amended) The isolated nucleic acid of Claim 1 having at least 90% nucleic acid sequence identity to:

(a) a nucleic acid sequence encoding the polypeptide of shown in Figure 88 (SEQ ID NO:88);

(b) a nucleic acid sequence encoding the polypeptide of shown in Figure 88 (SEQ ID NO:88), lacking its associated signal peptide;

(c) a nucleic acid sequence encoding the extracellular domain of the polypeptide of shown in Figure 88 (SEQ ID NO:88);

(d) a nucleic acid sequence encoding the extracellular domain of the polypeptide of shown in Figure 88 (SEQ ID NO:88) lacking its associated signal peptide;

(e) (c) the nucleic acid sequence of shown in Figure 87 (SEQ ID NO:87);

(f) (d) the full-length coding sequence of the nucleic acid sequence of shown in Figure 87 (SEQ ID NO:87); or

(g) (e) the full-length coding sequence of the cDNA deposited under ATCC accession number 203159;

wherein said isolated nucleic acid is more highly expressed in normal lung tissue compared to lung tumor, or wherein said isolated nucleic acid encodes a polypeptide that is more highly expressed in normal lung tissue compared to lung tumor.

4. (Currently Amended) The isolated nucleic acid of Claim 1 having at least 95% nucleic acid sequence identity to:

(a) a nucleic acid sequence encoding the polypeptide of shown in Figure 88 (SEQ ID NO:88);

(b) a nucleic acid sequence encoding the polypeptide of shown in Figure 88 (SEQ ID NO:88), lacking its associated signal peptide;

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(e) a nucleic acid sequence encoding the extracellular domain of the polypeptide of shown in Figure 88 (SEQ ID NO:88);

(d) a nucleic acid sequence encoding the extracellular domain of the polypeptide of shown in Figure 88 (SEQ ID NO:88) lacking its associated signal peptide;

(e) (c) the nucleic acid sequence of shown in Figure 87 (SEQ ID NO:87);

(f) (d) the full-length coding sequence of the nucleic acid sequence of shown in Figure 87 (SEQ ID NO:87); or

(g) (e) the full-length coding sequence of the cDNA deposited under ATCC accession number 203159;

wherein said isolated nucleic acid is more highly expressed in normal lung tissue compared to lung tumor, or wherein said isolated nucleic acid encodes a polypeptide that is more highly expressed in normal lung tissue compared to lung tumor.

5. (Currently Amended) The isolated nucleic acid of Claim 1 having at least 99% nucleic acid sequence identity to:

(a) a nucleic acid sequence encoding the polypeptide of shown in Figure 88 (SEQ ID NO:88);

(b) a nucleic acid sequence encoding the polypeptide of shown in Figure 88 (SEQ ID NO:88), lacking its associated signal peptide;

(c) a nucleic acid sequence encoding the extracellular domain of the polypeptide of shown in Figure 88 (SEQ ID NO:88);

(d) a nucleic acid sequence encoding the extracellular domain of the polypeptide of shown in Figure 88 (SEQ ID NO:88) lacking its associated signal peptide;

(e) (c) the nucleic acid sequence of shown in Figure 87 (SEQ ID NO:87);

(f) (d) the full-length coding sequence of the nucleic acid sequence of shown in Figure 87 (SEQ ID NO:87); or

(g) (e) the full-length coding sequence of the cDNA deposited under ATCC accession number 203159;

wherein said isolated nucleic acid is more highly expressed in normal lung tissue compared to lung tumor, or wherein said isolated nucleic acid encodes a polypeptide that is more highly expressed in normal lung tissue compared to lung tumor.

6. (Currently Amended) An isolated nucleic acid comprising:

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- (a) a nucleic acid sequence encoding the polypeptide of shown in Figure 88 (SEQ ID NO:88);
- (b) a nucleic acid sequence encoding the polypeptide of shown in Figure 88 (SEQ ID NO:88), lacking its associated signal peptide;
- (c) a nucleic acid sequence encoding the extracellular domain of the polypeptide of shown in Figure 88 (SEQ ID NO:88);
- (d) a nucleic acid sequence encoding the extracellular domain of the polypeptide of shown in Figure 88 (SEQ ID NO:88) lacking its associated signal peptide;
- (e) (c) the nucleic acid sequence of shown in Figure 87 (SEQ ID NO:87);
- (f) (d) the full-length coding sequence of the nucleic acid sequence of shown in Figure 87 (SEQ ID NO:87); or
- (g) (e) the full-length coding sequence of the cDNA deposited under ATCC accession number 203159.
7. (Currently Amended) The isolated nucleic acid of Claim 6 comprising a nucleic acid sequence encoding the polypeptide of shown in Figure 88 (SEQ ID NO:88).
8. (Currently Amended) The isolated nucleic acid of Claim 6 comprising a nucleic acid sequence encoding the polypeptide of shown in Figure 88 (SEQ ID NO:88), lacking its associated signal peptide.
9. (Cancelled)
10. (Cancelled)
11. (Currently Amended) The isolated nucleic acid of Claim 6 comprising the nucleic acid sequence of shown in Figure 87 (SEQ ID NO:87).
12. (Currently Amended). The isolated nucleic acid of Claim 6 comprising the full-length coding sequence of the nucleic acid sequence of shown in Figure 87 (SEQ ID NO:87).
13. (Original) The isolated nucleic acid of Claim 6 comprising the full-length coding sequence of the cDNA deposited under ATCC accession number 203159.
14. (Currently Amended) An isolated nucleic acid that hybridizes under stringent conditions to:
- (a) a nucleic acid sequence encoding the polypeptide of shown in Figure 88 (SEQ ID NO:88);

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- (b) a nucleic acid sequence encoding the polypeptide of shown in Figure 88 (SEQ ID NO:88), lacking its associated signal peptide;
- (c) a nucleic acid sequence encoding the extracellular domain of the polypeptide of shown in Figure 88 (SEQ ID NO:88);
- (d) a nucleic acid sequence encoding the extracellular domain of the polypeptide of shown in Figure 88 (SEQ ID NO:88) lacking its associated signal peptide;
- (e) (c) the nucleic acid sequence of shown in Figure 87 (SEQ ID NO:87);
- (f) (d) the full-length coding sequence of the nucleic acid sequence of shown in Figure 87 (SEQ ID NO:87); or
- (g) (e) the full-length coding sequence of the cDNA deposited under ATCC accession number 203159;

wherein said stringent conditions comprise 50% formamide, 5 x SSC (0.75 M NaCl, 0.075 M sodium citrate), 50 mM sodium phosphate (pH 6.8), 0.1% sodium pyrophosphate, 5 x Denhardt's solution, sonicated salmon sperm DNA (50 µg/ml), 0.1% SDS, and 10% dextran sulfate at 42°C, with washes at 42°C in 0.2 x SSC (sodium chloride/sodium citrate) and 50% formamide at 55°C, followed by a high-stringency wash consisting of 0.1 x SSC containing EDTA at 55°C.

15. (Canceled).

16. (Original) The isolated nucleic acid of Claim 14 which is at least 10 nucleotides in length.

17. (Original) A vector comprising the nucleic acid of Claim 1.

18. (Original) The vector of Claim 17, wherein said nucleic acid is operably linked to control sequences recognized by a host cell transformed with the vector.

19. (Currently Amended) A An isolated host cell comprising the vector of Claim 17.

20. (Original) The host cell of Claim 19, wherein said cell is a CHO cell, an E. coli or a yeast cell.

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DELETION OF INVENTORS

Please correct the inventorship under 37 CFR §1.48(b) by removing the following inventors from the present application:

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